

FIGURE 1A

1 CCCC GCGTCGGTCTTCCACCTCACCTTTTCGAGCTGGCCGCCGCTTGCTGTGCGCAGTTTC 60  
61 GGGGGACTGGACCTTCCCTGGCTTTTAGCAGCGCCGAGCGCCATGGCGACCCTTTGCTGG 120  
121 GCAGGTGACCGATTCCGGGTGCCCGAAGGAGCTGGCGTGGGTCTGCCTTGCAGCCGCCCG 180  
181 CCTGGACAGGATGTTTGTAGAGGGCTGAAGAGGAAATATGGTGACCAGGAAGAAGGAGT 240  
1 M F A R G L K R K Y G D Q E E G V 17  
241 AGAGGGTTTTGGCACTGTCCCTTCTATAGCCTGCAGCGACAGTCACTCCTGGACATGTC 300  
E G F G T V P S Y S L Q R Q S L L D M S 37  
301 CCTTGTCAAGCTCCAGCTCTGTACATGCTAGTGGAGCCCAATCTCTGCCGCTCGGTCT 360  
L V K L Q L C H M L V E P N L C R S V L 57  
361 CATCGCCAACACAGTCCGGCAGATCCAGGAGGAAATGAGCCAGGATGGTGTGTGGCATGG 420  
I A N T V R Q I Q E E M S Q D G V W H G 77  
421 GATGGCACCCCAGAATGTAGATCGGGCACCAGTTGAACGCCTGGTGTCCACAGAGATCCT 480  
M A P Q N V D R A P V E R L V S T E I L 97  
481 GTGTCGTACAGTGAGGGGAGCTGAGGAAGAGCACCCTGCTCCTGAACTGGAAGATGCTCC 540  
C R T V R G A E E E H P A P E L E D A P 117  
541 CTTGCAAACTCGGTTTCCGAGCTCCCCATCGTTGGCTCAGCACCAGGGCAAAGGAACCC 600  
L Q N S V S E L P I V G S A P G Q R N P 137  
601 TCAGAGCAGCCTCTGGGAGATGGACAGCCCAAGAAAACAGGGGAAGCTTTTCAGAAGTC 660  
Q S S L W E M D S P Q E N R G S F Q K S 157  
661 ACTGGACCAGATATTTGAGACCCTGGAGAACAACAACTCCAGTTCACTGGAGGAACCTCTT 720  
L D Q I F E T L E N K N S S S V E E L F 177  
721 CTCAGATGTGGACAGCTCCTACTATGACCTGGACACAGTGCTAACAGGAATGATGAGTGG 780  
S D V D S S Y Y D L D T V L T G M M S G 197  
781 GACCAAGTCCAGTCTCTGCAATGGCCTTGAGGGCTTTGCTGCAGCCACCCCTCCTCCCAG 840  
T K S S L C N G L E G F A A A T P P S 217  
841 TTCCACTTGCAAGTCTGGCTGAGCTGAGCATGTGGTAGAGATTCTGGTGGAGAC 900  
S T C K S D L A E L D H V V E I L V E T 237  
901 CTGAGAGGCCACCCAGTGGGCTAAGGGTGAGGCCACCAGTCCCCATGGAGCTCACGTGT 960  
\*  
961 GTTGTGACCCAGAGACAGATAAGCACTTGTCTTAAGAGGGGCTCTGGCTCTTGAGCTCAT 1020  
1021 TATCCTTTTGTGTGACATTGGACTCACTGTGGAGGATGGTGTGTACAGCTATGTCTAGT 1080  
1081 CTATTTTCAATTAGATAGCTGAACCTTTCTAAAATTAAGTTTATATGTTTTTGGGCAATA 1140  
1141 TTTTGTCTTAAGATATATTTTAAACTTTTTTATACTTTAGATTTTTTTCAGCTATTTTC 1200  
1201 TTAAAAGTATATTTTTTCTACAAACATCCTCTGCTGCTACATTAGAAACATTTATAACCT 1260  
1261 AAATACGATTGGTGTGTCAATTTTAAAGGTTTAAATAGAAAACCTTCTTTTGTACTGAGTC 1320  
1321 TCTACACTCCCAAGGCAACTGTAAATGTAGCCGGCCGGGTGTTTACATGAGAGGCTCCAG 1380  
1381 TATGGTCTACATTCTAGTAGAGCTTGAAAAGAACCATGCACAGCTCCACTGCCCCCTCAC 1440  
1441 TGGGTCTGCTCTGGCGGATCGGAGCTCTCTTCTAGCCCCGTGTGCAGGATGGCTTTATT 1500  
1501 TATGCCTATTTATATGTAAATGCCACTGAAAGCTAAGGTCTTACTCCTGGAAATCCCAAC 1560  
1561 ACCAGTTCTTCAGGGACTGCTGTGAGGCAGTGCCTTATGCAGGTCTTGTCTTTGGCCATC 1620  
1621 ACTGTCTGGTTCCAGCCAGCACATGTGACATGAGGACATGACATGCCCCGAACCACCCA 1680  
1681 GCACCACATGCTCCATGTCAAGTGTGTACGTGGAGACCACTGGCTCCCAGGCCTGTGCTC 1740  
1741 AGAGAGGGTGTGCAGTCTACGTGTGCTGGGGGGACGACGGTGACCTGTGCTTGCTTGC 1800  
1801 TTTTAAAATGGTGTCTGGACGTTTAAAGGTTAAAAACAATCCGACTCCATATGATTTAGG 1860  
1861 GCTCCTCCACCCTGGGGTGGCCCTATGCTGTCTGCTTGGATCTCAAAGTCTTGGTACTC 1920  
1921 GGCAGTGTGAGACTCCACCCATGTATCCTTTTTGTTTCTTGTGCTTTTTTTGGACTT 1980  
1981 CCCAACCTGAGCCTAAGGTTTATTTTATATGTGCTTCAATATCAACAATGTAAACCTCA 2040  
2041 CTTTATTAAAGTATCCAGCAAATGGAAAAA

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FIGURE 1B

1 GGAAGCTGGCGGCACAGCCGTGGCGCCTGGCTGAGCAGAGGACCCGGCGGGCGGCCTCG 60  
61 CGGGTCAGGACACAATGTTTGCACGAGGACTGAAGAGGAAATGTGTTGGCCACGAGGAAG 120  
1 M F A R G L K R K C V G H E E D 16  
121 ACGTGGAGGGAGCCCTGGCCGGCTTGAAGACAGTGTCTCATACAGCCTGCAGCGGCAGT 180  
V E G A L A G L K T V S S Y S L Q R Q S 36  
181 CGCTCCTGGACATGTCTCTGGTGAAGTTGCAGCTTTGCCACATGCTTGTGGAGCCCAACC 240  
L L D M S L V K L Q L C H M L V E P N L 56  
241 TGTGCCGCTCAGTCCCTCATTGCCAACACGGTCCGGCAGATCCAAGAGGAGATGACGCAGG 300  
C R S V L I A N T V R Q I Q E E M T Q D 76  
301 ATGGGACGTGGCGCACAGTGGCAGACCCCGGCTGCAGAGCGGGCGCCGCTCGACCGCTTGG 360  
G T W R T V A P Q A A E R A P L D R L V 96  
361 TCTCCACGGAGATCCTGTGCGGTGCAGCGTGGGGGCAAGAGGGGGCACATCCTGCTCCTG 420  
S T E I L C R A A W G Q E G A H P A P G 116  
421 GCTTGGGGGACGGCCACACAGGGTCCAGTTTCTGACCTTTGCCAGTCACCTCAGCAC 480  
L G D G H T Q G P V S D L C P V T S A Q 136  
481 AGGCACCAAGGCACCTGCAGAGCAGCGCCTGGGAGATGGATGGCCCTCGAGAAAACAGAG 540  
A P R H L Q S S A W E M D G P R E N R G 156  
541 GAAGCTTTCACAAGTCACCTTGATCAGATATTTGAAACGCTGGAGACTAAAAACCCAGCT 600  
S F H K S L D Q I F E T L E T K N P S C 176  
601 GCATGGAAGAGCTGTTCTCAGACGTGGACAGCCCTACTACGACCTGGACACAGTACTGA 660  
M E E L F S D V D S P Y Y D L D T V L T 196  
661 CAGGCATGATGGGGGTGCCAGGCCGGGCCCCCTGCCAAGGGCTCGAGGGCTTGGCTCCGG 720  
G M M G G A R P G P C E G L E G L A P A 216  
721 CCACCCAGGCCCTCAGCTGCAAGTCCGACCTGGGCGAGCTGGACCACGTGGTGG 780  
T P G P S S S C K S D L G E L D H V V E 236  
781 AGATCCTGGTGGAGACCTGAGCAGGAGCCCTGAGTGTCTACAGCCGCCTCTGACGCATTG 840  
I L V E T \* 241  
841 ACACGTGAGCACTGGCTCCACGAGGGTGCAGCTGCCGCCAGCGGCCAGCCTTGCTGC 900  
901 CCTGTCTGCTGATTCTGAGAAATCCCAGAACAGCCATTACCAGTGGGGCTGCAGCCCTA 960  
961 GGCCCGTCCCACTCACCTCCCCCTGTGGAGCGCCAGGCAGAGGCTGTTCTGGAAGGCTT 1020  
1021 CTTGTCTTCTGACGTCCCCACAGCCCTGGGCCCCCTCGTGTCTCTTTGTGTCCCCCACTGT 1080  
1081 AGAGGACGGTGAGCCGCAGCTGCATCAACCTCCTTTTACCTTTAGATAGGTGAATTTTAA 1140  
1141 CAATTCAGTTTTACATGTTTTGGGCAGTATTTTGTCTTAAGATATATTTTTTAACTTTT 1200  
1201 TATACCTTATCTCTTTAGATTTTTTTCAGCTATTTTCTTAAAAGTATATTTTTTCTATAAA 1260  
1261 CATCCTTTGCTGCTACATTAGAATTTTTATAGCCTAAACAATTGCAGTTGGTGTGTTTCA 1320  
1321 TTTTTTTAAGGTTTAAATAAGGGTTTTTTGTTTTGTTTTGTTTTTGCAGTGAGCATCAC 1380  
1381 TACAGTCTCAGTCAACAGTGTGAATGTATCATGTTTTACTTTAAATGTGTGTGTGATACT 1440  
1441 TCTTCATTATGTCCTGCGCTGCAGTGAGACCTGGGTGAAATCAGGAGCCGCACACAGCC 1500  
1501 ACATCTTCCTAGACCTAAGAGTAAATTATGGAGGATTTTATTTATGTCTATTTATATGTA 1560  
1561 AATGTCAATTGAAGACAAAGGTCAAATATTTGTCTGTTTGTAGATCACAGGCACCAAGTTGG 1620  
1621 TCTTCAGGGACCTCATAGCCCCCTCGGTGGTGCCTTCTCAAGGCAGTGTTCCTGGAGGCTC 1680  
1681 CCATCAGGGTCAGCCCATGCACCTGCCCTGGGTGAGGAAGTAGCATTGCTGCTGGATGAG 1740  
1741 AAACGCCTGCGCTGCTCTGTAGACTGGTGTGAAACAAAAGGTTAAGGCTAGGTTGAAG 1800  
1801 TCTAGAATGAAAGAAATCTGAATCCATGTCAATTCATAACCCCTTGATCTGTAGTGTCTATG 1860  
1861 GGTGCTGCCGCAGGCAGGGAGTGAGCTGGGGGTGCCTGCAGCCTTCCACTCCTGCCCCGC 1920  
1921 CTCACCCACATGCTCCCTGTTTCTCATGCTTTCTCTAACTTCCTCACCCTTAACCAAAA 1980  
1981 AAGGTGTGTTTTCTTTTGTGCATATAGCCATTCTTAAATATCAGTGATGTAAACCTCACT 2040  
2041 TTATTAATAAAATATCCAGCAAAAAAAAAAAAAAAAAAAAAA

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FIGURE 2

Mouse <i>Hepp</i>	1	MFARGLKRKYG----DOEEGVGFGTVPSYSLQRQSLDMSLVKLQLCHMLVEPNLCRSV
Human <i>HEPP</i>	1	MFARGLKRKCVGHEEDVEGALAGLKTSSYSLQRQSLDMSLVKLQLCHMLVEPNLCRSV
Mouse <i>Hepp</i>	57	LIANTVROIQEEMSQDGVWEGMAPONVDRAPVERLVSTEILCRTVRGAE EHPAPELEDA
Human <i>HEPP</i>	61	LIANTVROIQEEMTQDGTWRTVAPOAAERAPLDRLVSTEILCRAAWGOEGAHAPAGLGDG
Mouse <i>Hepp</i>	117	PLQNSVSELPVIGSAPGORNPOSSLWEMDSPQENRGSFQKSLDQIFETLENKNSSSVEEL
Human <i>HEPP</i>	121	HTQGPVSDLCPVTSAGAPRHLOSSAWEMDGPRENRGSFHKSLDQIFETLETKNPSCEEL
Mouse <i>Hepp</i>	177	FSDVDSSYYDLDTVLTGMMSCGKSSLCNGLECFAAATPPSSCKSDLAELDHVVEILVE
Human <i>HEPP</i>	181	FSDVDSPYYDLDTVLTGMMGCAEPGPCEGLEGLAPATPGPSSCKSDLGELDHVVEILVE
Mouse <i>Hepp</i>	237	T
Human <i>HEPP</i>	241	T

FIGURE 3

Zebrafish <i>Hepp</i>	1	MFSKGLKRKFADGGEIISDGLVAARVASSYSLQRQSLDMSLVKLQLCHMLVEPNLCRS
Mouse <i>Hepp</i>	1	MFARGLKRKYG----DOEEGVGFGTVPSYSLQRQSLDMSLVKLQLCHMLVEPNLCRS
Human <i>HEPP</i>	1	MFARGLKRKCVGH-EEDVEGALAGLKTSSYSLQRQSLDMSLVKLQLCHMLVEPNLCRS
Zebrafish <i>Hepp</i>	61	VLIANTVROIQEEMTHDGSWEMVTEAFCGASQSPSERLVATEVLCR-----
Mouse <i>Hepp</i>	56	VLIANTVROIQEEMSQDGVWEGMAPONVDR--APVERLVSTEILCRTVRGAE EHPAPEL
Human <i>HEPP</i>	60	VLIANTVROIQEEMTQDGTWRTVAPOAAER--APLRLVSTEILCRAAWGOEGAHAPAGL

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FIGURE 4B

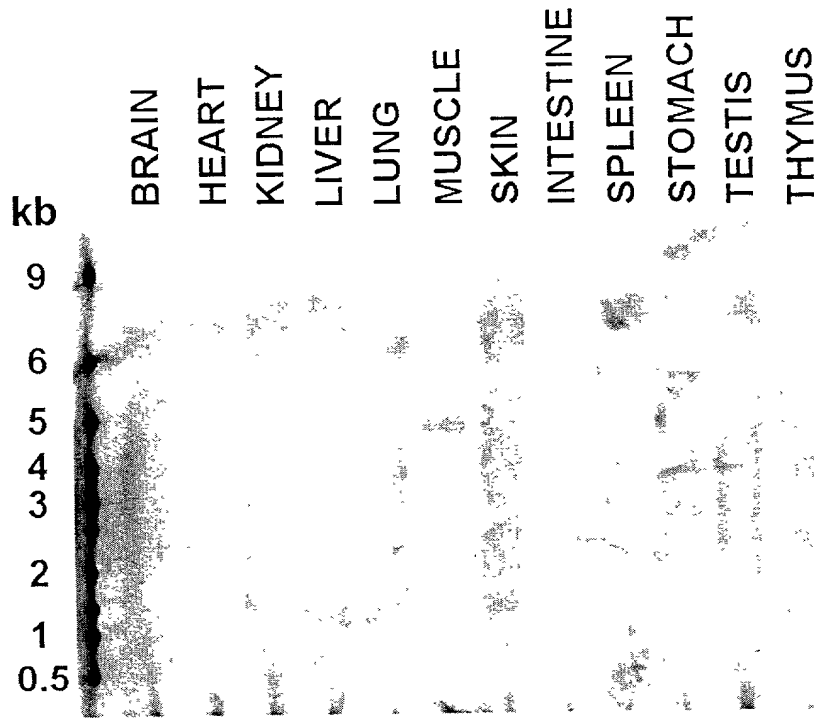


FIGURE 4A

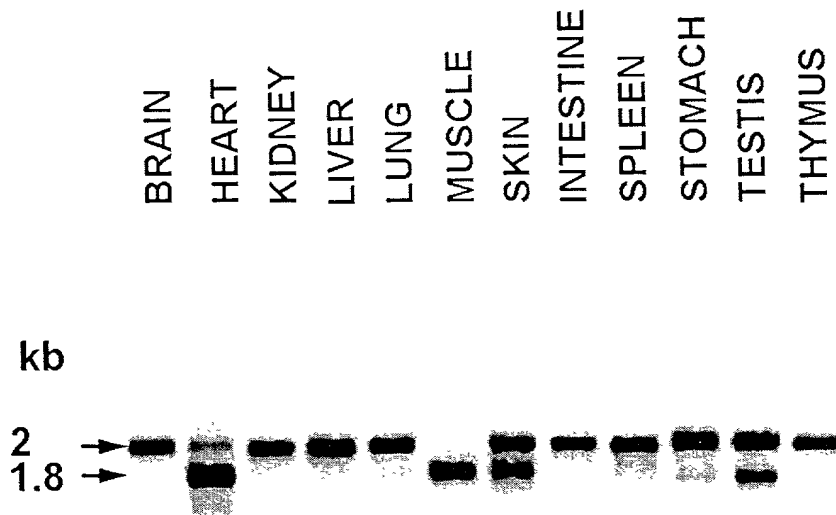
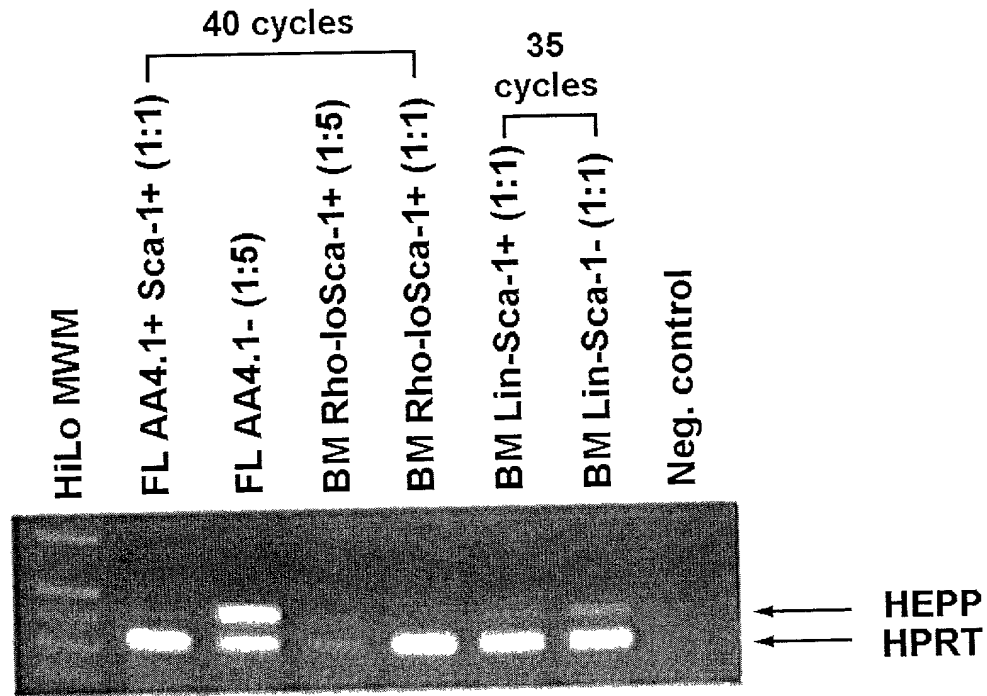


FIGURE 5



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 FIGURE 6

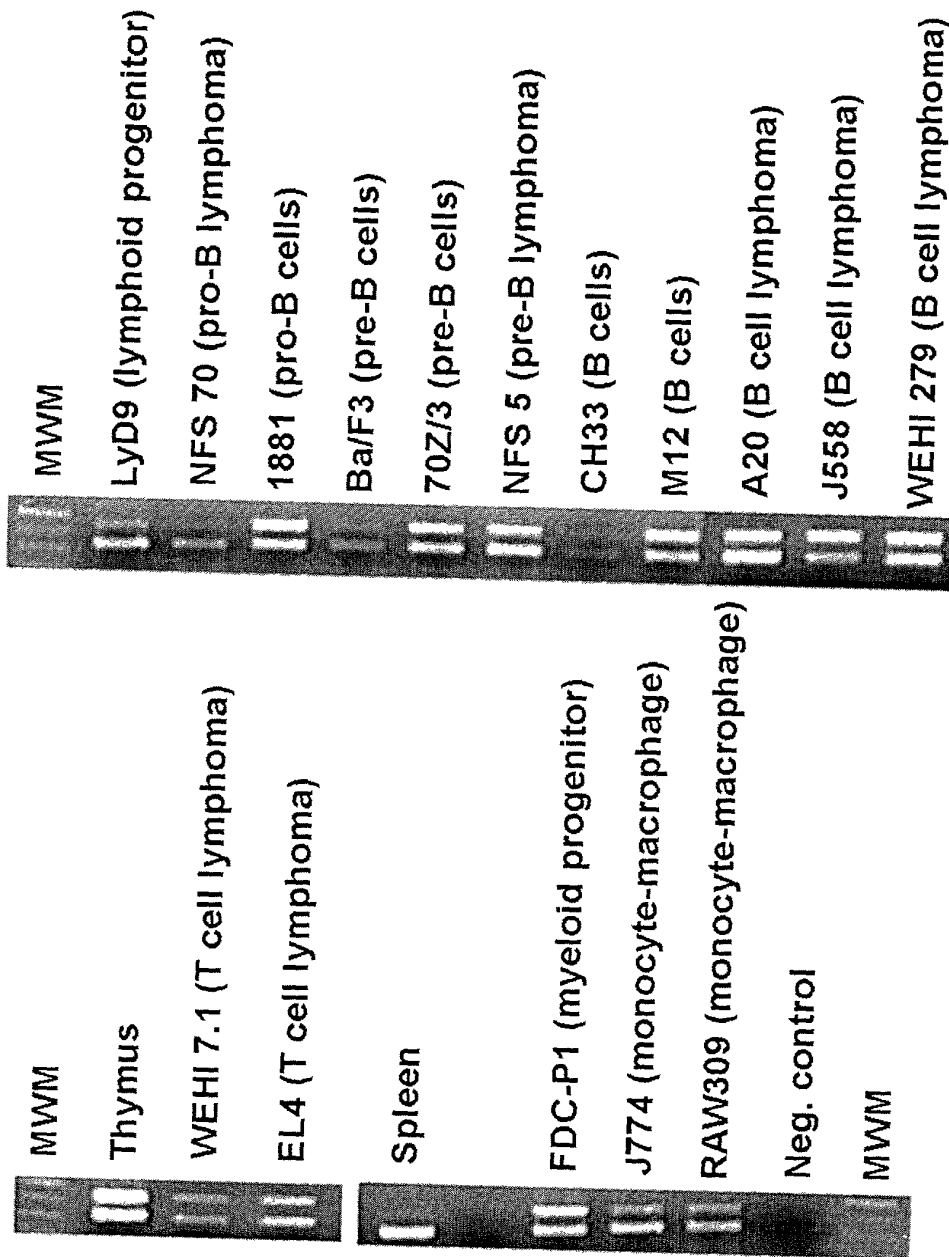
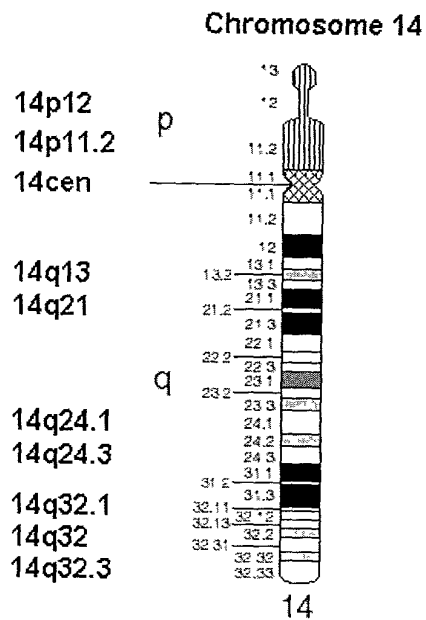


FIGURE 7



translocation breakpoints

FIGURE 8

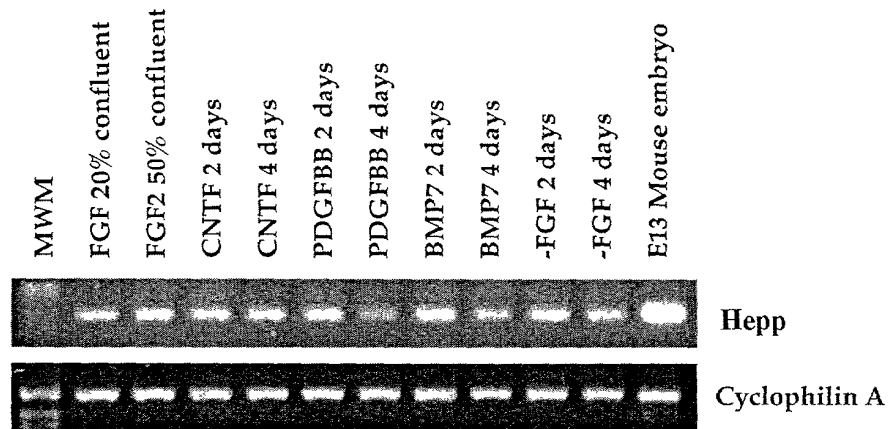
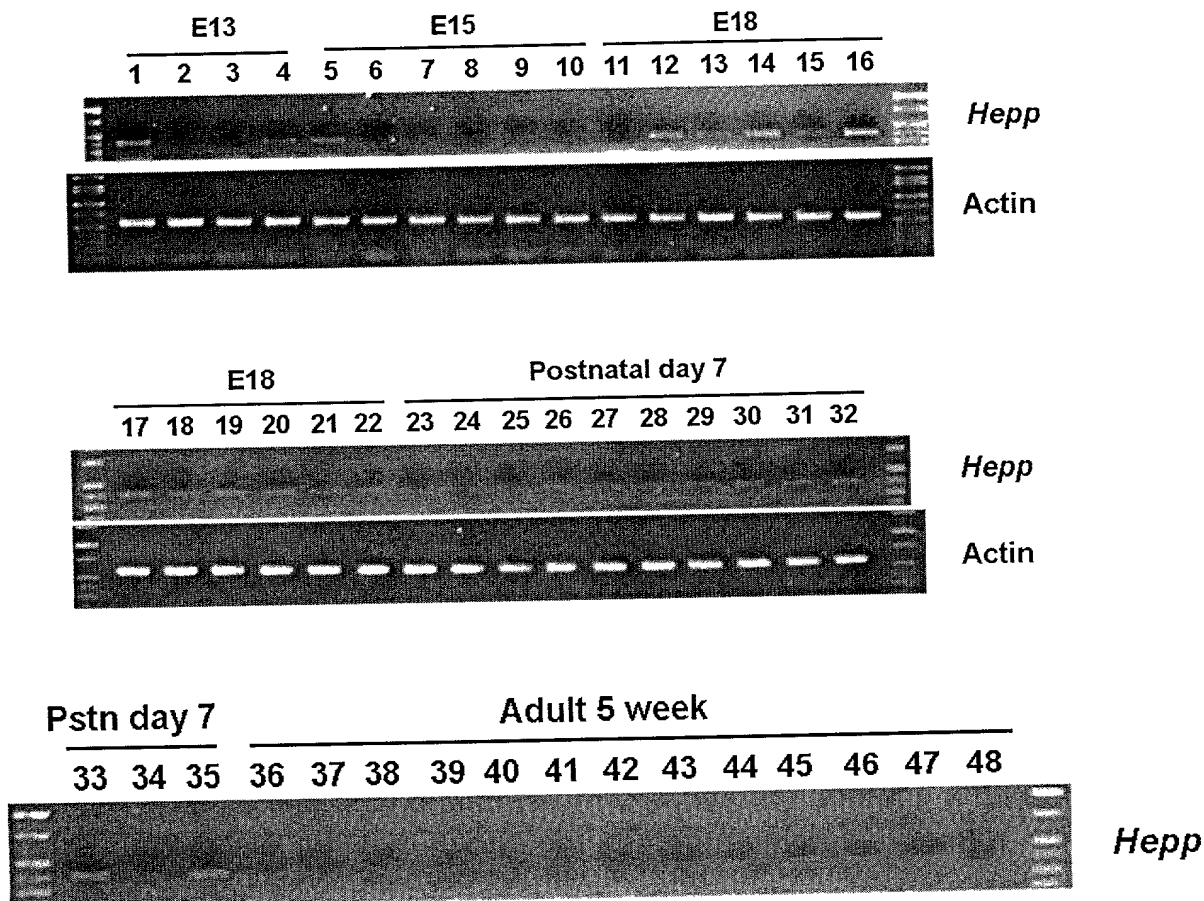




FIGURE 9



Embryo day 13	Embryo day 18	Postnatal day 7	Adult 5 week
1. Telencephalon/Diencephalon	11. Frontal cortex	23. Frontal cortex	36. Frontal cortex
2. Mesencephalon (Midbrain)	12. Posterior cortex	24. Posterior cortex	37. Posterior cortex
3. Rhombencephalon (Hindbrain)	13. Entorhinal cortex	25. Entorhinal cortex	38. Entorhinal cortex
4. Spinal cord	14. Olfactory bulb	26. Olfactory bulb	39. Olfactory bulb
	15. Hippocampus	27. Hippocampus	40. Hippocampus
	16. Striatum	28. Striatum	41. Striatum
	17. Thalamus	29. Thalamus	42. Thalamus
	18. Hypothalamus	30. Hypothalamus	43. Hypothalamus
	19. Midbrain	31. Cerebellum	44. Cerebellum
	20. Pons	32. Midbrain	45. Midbrain
	21. Medulla	33. Pons	46. Pons
	22. Spinal cord	34. Medulla	47. Medulla
		35. Spinal cord	48. Spinal cord

FIGURE 10

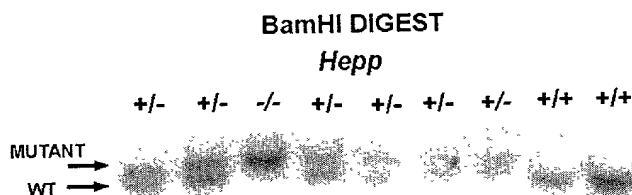
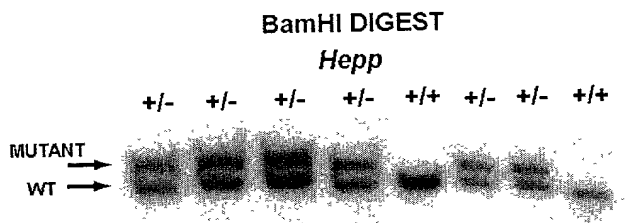


FIGURE 11

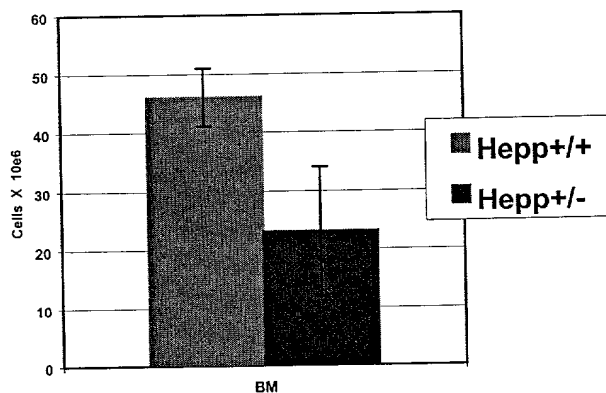


FIGURE 12

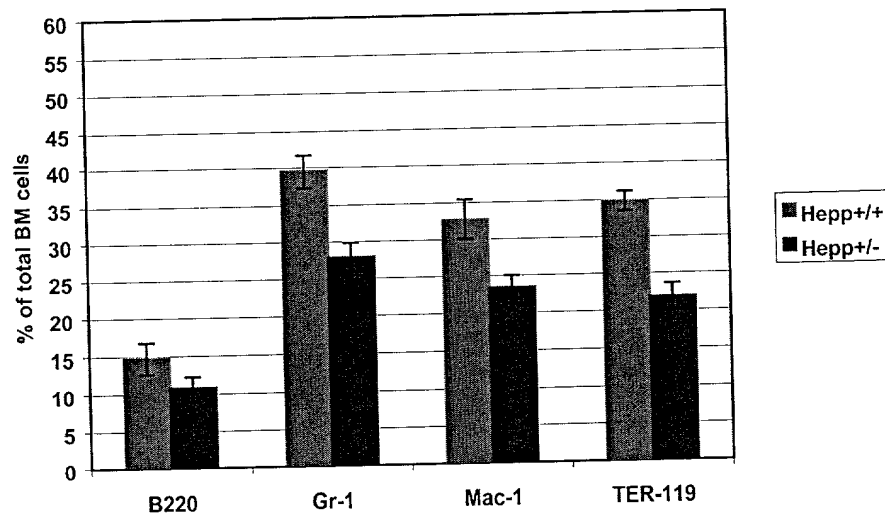


FIGURE 13

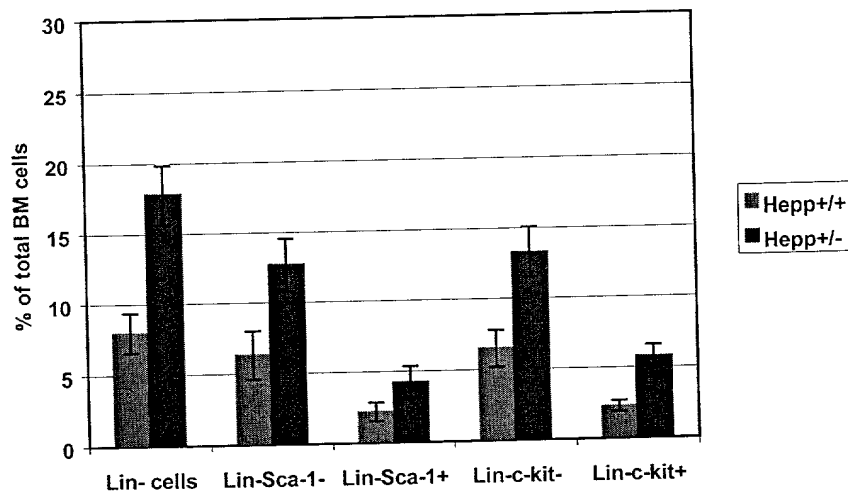


FIGURE 14

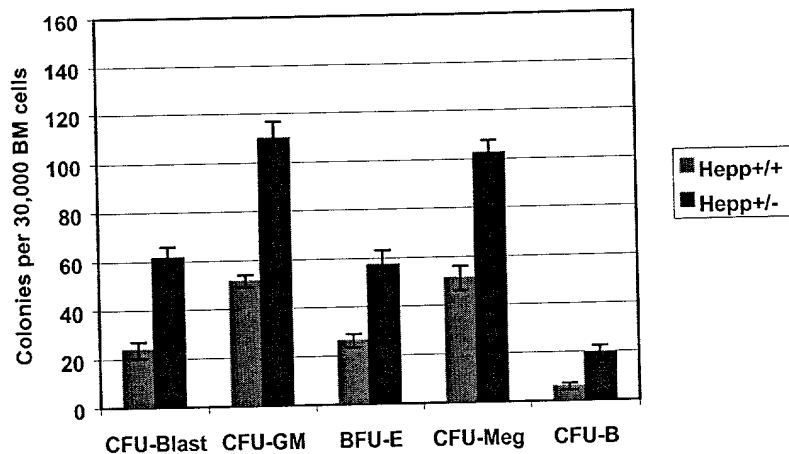


FIGURE 15A-B

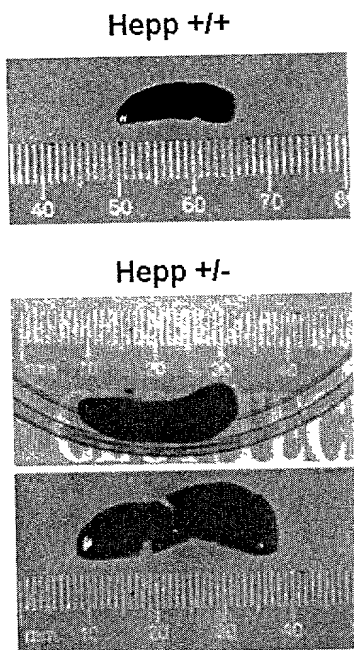


FIGURE 15C

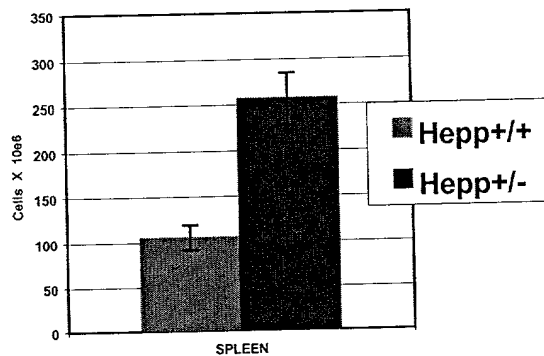


FIGURE 16

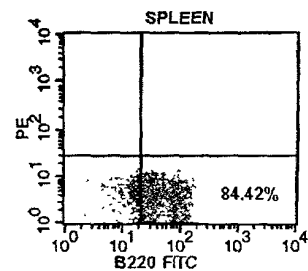
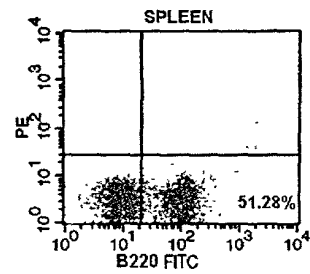
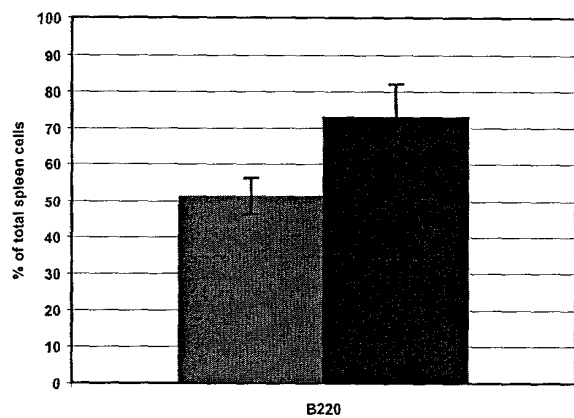


FIGURE 17

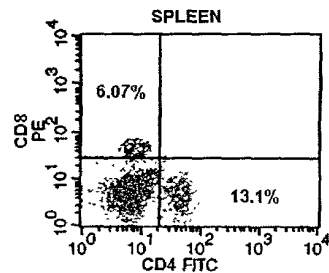
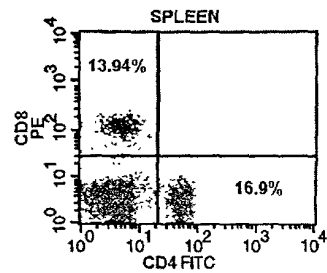
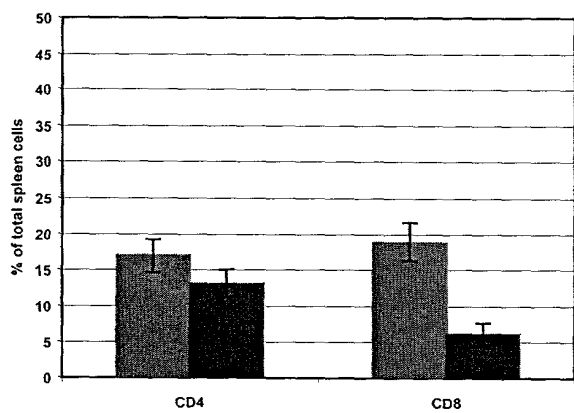


FIGURE 18A

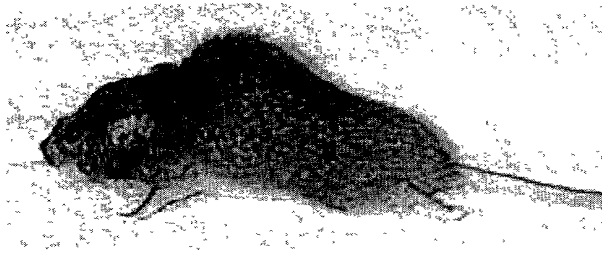


FIGURE 18B

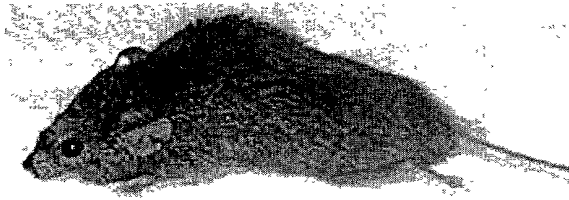
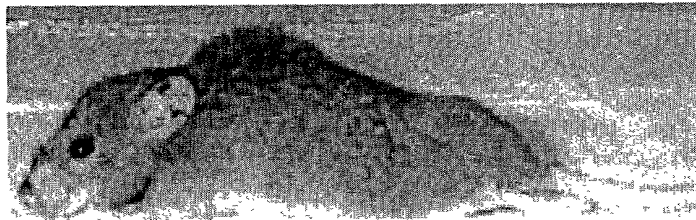


FIGURE 18C



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